Ensemble Learning

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IIT Gandhinagar

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Outline

1. Introduction to Ensemble Learning

2. Why Ensembles Work: Theoretical Foundation

Core Idea

Ensemble Learning: Combine multiple weak learners to create a strong learner

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Everyday Analogy

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Key Insight:

Ensemble Performance > Individual Model Performance

Pop Quiz: Ensemble Intuition

Quick Quiz 1

Why might combining multiple models work better than using a single model?

a) More models always mean better performance

Answer: b) When models make different errors, their combination can correct individual mistakes!

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Why might combining multiple models work better than using a single model?

- a) More models always mean better performance
- b) Different models make different types of errors
- c) Ensemble models are faster to train

Answer: b) When models make different errors, their combination can correct individual mistakes!

Based on Ensemble methods in ML by Dietterich

Three reasons why ensembles make sense:

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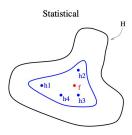
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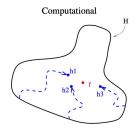
E.g., we can learn many decision trees for the same data giving the same accuracy.



2) Computational: Even if data is sufficient, some classifiers/regressors can get stuck in local optima or apply greedy strategies. Computationally learning the "best" hypothesis can be non-trivial.

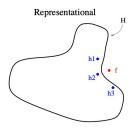
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E.g., decision trees employ greedy criteria



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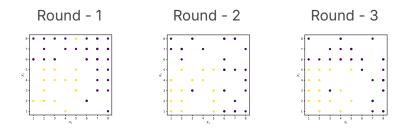
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- 3) Two classifiers are diverse: if they make different errors on new data points

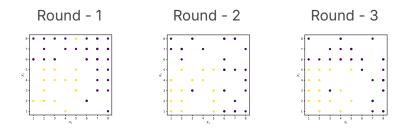
If the three classifiers are identical, i.e. not diverse, then when $h_1(x)$ is wrong $h_2(x)$ and $h_3(x)$ will also be wrong.

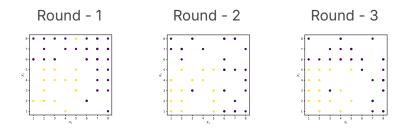
If the three classifiers are identical, i.e. not diverse, then when $h_1(x)$ is wrong $h_2(x)$ and $h_3(x)$ will also be wrong. However, if the errors made by the classifiers are uncorrelated, then when $h_1(x)$ is wrong, $h_2(x)$ and $h_3(x)$ may be correct, so that a majority vote will correctly classify.

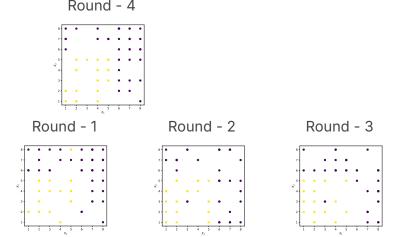
Error Probability of each model = ε = 0.3

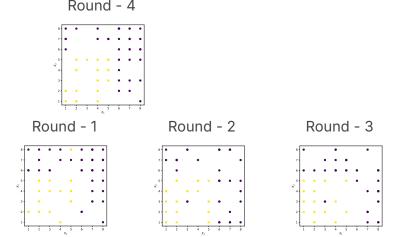
$$\begin{aligned} \textit{Pr}(\text{ensemble being wrong}) &= \\ {}^3\textit{C}_2(\varepsilon^2)(1-\varepsilon)^{3-2} + {}^3\textit{C}_3(\varepsilon^3)(1-\varepsilon)^{3-3} \\ &= 0.19 < 0.3 \end{aligned}$$

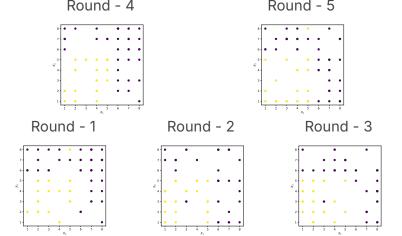


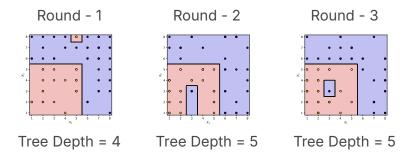


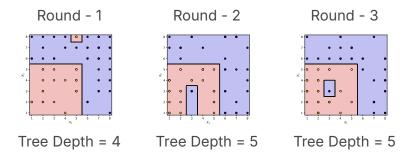


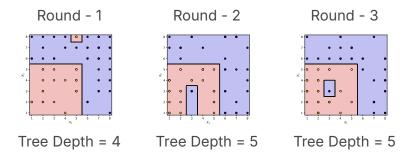


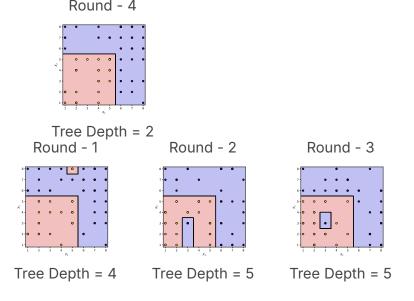


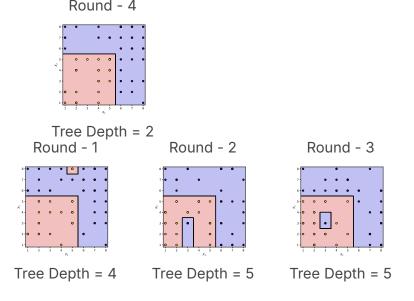


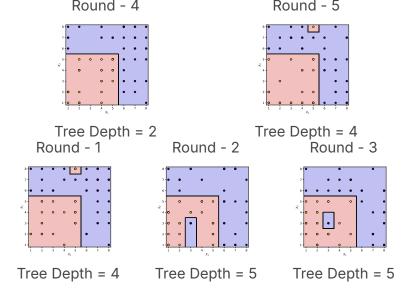










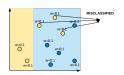


All learners are incrementally built.

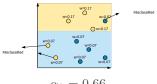
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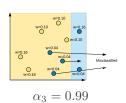
Incremental building: Incrementally try to classify "harder" samples correctly.

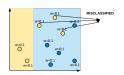


 $\alpha_1 = 0.42$

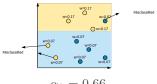




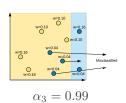


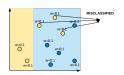


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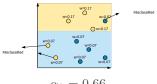




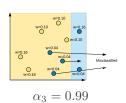


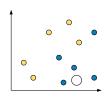


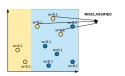
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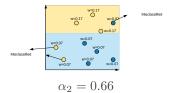


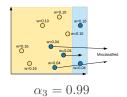


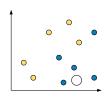


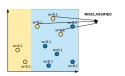


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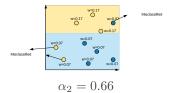


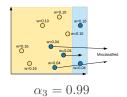


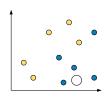


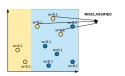


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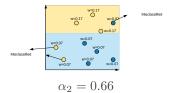


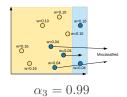






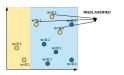
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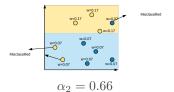




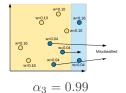


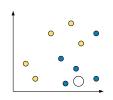


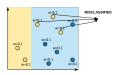
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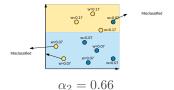
Let us say, yellow class is +1 and blue class is -1



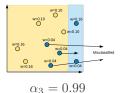




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Let us say, yellow class is +1 and blue class is -1 Prediction = SIGN(0.42*-1 + 0.66*-1 + 0.99*+1) = Negative = blue



Notebook: boosting-explanation.html

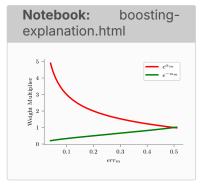
0.2 0.3

 err_m

0.4 0.5

0.0

0.1



ADABoost for regresion

From Paper: Improving Regressors using Boosting Techniques

Our problem will be that the modeling error is also nonzero because we have to determine the model in the presence of noise. Since we don't know the probability distributions, we approximate the expectation of the ME and PE using the sample ME (if the truth is known) and sample PE and then average over multiple experiments.

In the following discussion, we detail both bagging and boosting. We then discuss how to build trees which are the basic building blocks of our regression machines and use these ensembles on some standard test functions.

2. BAGGING

The following is a paraphrase of Breiman (1996b) with some difference in notation. Suppose we pick with replacement N1 examples from the training set of size N_{+} and call the k'th set of observations O_{+} . Based on these observations, we form a predictor $v^{(p)}(\mathbf{x}, O_1)$. Because we are sampling with replacement, we may have multiple observations or no observations of a particular training example. Sampling with replacement is sometimes termed bootstrap sampling [Efron and Tibshirani (1993)] and therefore this method is called bootstrap aggregating or bagging for short. The ensemble predictor is formed from the approximation to the expectation over all the observation sets, i.e. $E_O[y^{(p)}(x,O)]$ by using the average of the outputs of all the predictors. Breiman discusses which algorithms are good candidates for predictors and concludes that the best predictors are unstable, i.e., a small change in the training set O_k causes a large change in the predictor $v^{(p)}(\mathbf{x}, O_k)$. Good candidates are regression trees and neural nets.

3. BOOSTING

In bagging, each training example is equally likely to be picked. In boosting, the probability of a particular example being in the training set of a particular machine depends on the performance of the prior machines on that example. The following is a modification of Adaboost R [Freund and Schapire (1996a)].

Initially, to each training pattern we assign a weight $w_i=1$ $i=1,...,N_1$

Repeat the following while the average loss \overline{L} defined

set. Each machine makes a hypothesis: h.:x→v

3. Pass every member of the training set through this machine to obtain a prediction $v_i^{(p)}(x_i)$ $i=1,...,N_1$.

4. Calculate a loss for each training sample $L_l = L \left[|y_j^{(p)}(\mathbf{x}_l) - y_l| \right]$. The loss L may be of any functional form as long as $L \in [0,1]$. If we let

 $D=\sup |y_i^{(p)}(\mathbf{x}_i) - y_i|$ $i=1,...,N_1$ then we have three candidate loss functions:

$$L_i = \frac{\mid y_i^{(p)}(x_i) - y_i \mid}{D} \quad (linear)$$

$$L_i = \frac{\mid y_i^{(p)}(\pmb{x}_i) - y_i \mid^2}{D^2} \qquad (square\ law)$$

$$L_{i} = 1 - exp \left[\frac{- \mid y_{i}^{(p)}(\mathbf{x}_{i}) - y_{i} \mid}{D} \right] \qquad (exponential)$$

5. Calculate an average loss: $\bar{L} = \sum_{i=1}^{N_1} L_i p_i$

6. Form $\beta = \frac{\overline{L}}{1 - \overline{L}}$. β is a measure of confidence in the predictor. Low β means high confidence in the prediction.

7. Update the weights: w₁→w₁B**[1-L₁], where ** indicates exponentiation. The smaller the loss, the more the weight is reduced making the probability smaller that this pattern will be picked as a member of the training set for the next machine in the ensemble.

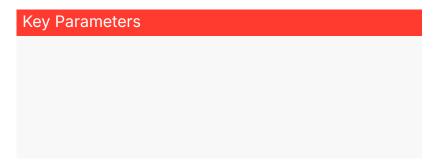
 For a particular input x_i, each of the T machines makes a prediction h_t, t=1,...,T. Obtain the cumulative prediction h_t using the T predictors:

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- As features are randomly selected, we learn decorrelated trees and helps in reducing variance.



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Training Algorithm

For each tree $i \in \{1, \dots, N\}$:

- 1. **Bootstrap sampling:** Select *n* samples with replacement
- 2. Train decision tree: Learn on bootstrap sample
- 3. **Random feature selection:** At each split, consider only *m* random features

Pop Quiz: Random Forest

Quick Quiz 3

What happens if we set m = M (use all features) in Random Forest?

a) Trees become more diverse

Answer: b) Using all features makes trees more similar, reducing diversity!

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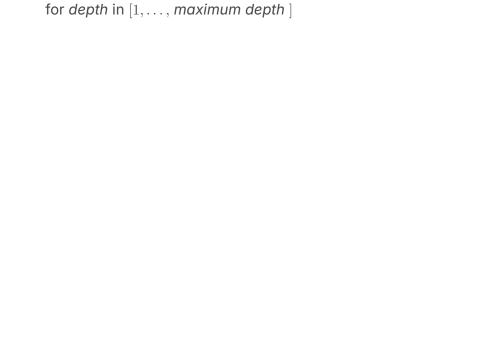
- a) Trees become more diverse
- b) Trees become more correlated
- c) No effect on performance

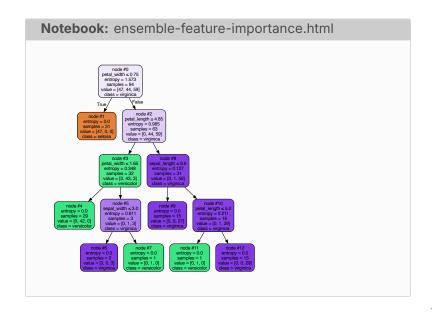
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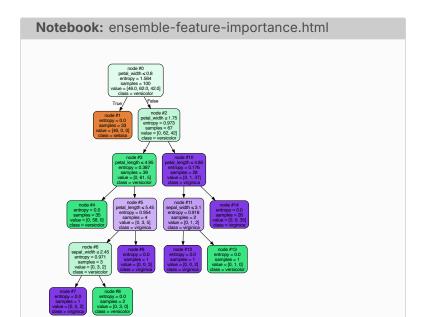
Dataset

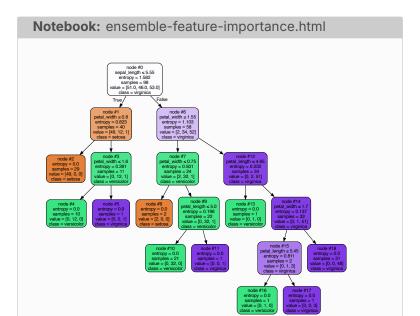
	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa
145	6.7	3.0	5.2	2.3	virginica
146	6.3	2.5	5.0	1.9	virginica
147	6.5	3.0	5.2	2.0	virginica
148	6.2	3.4	5.4	2.3	virginica
149	5.9	3.0	5.1	1.8	virginica

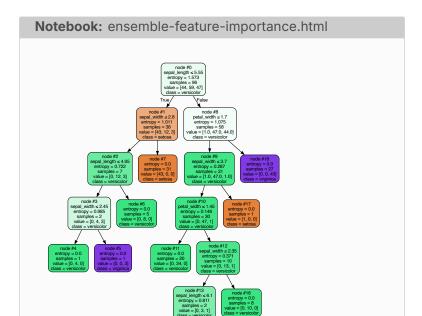
150 rows × 5 columns

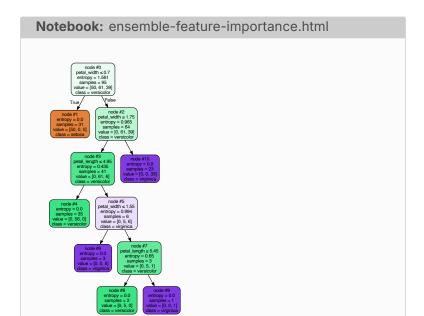


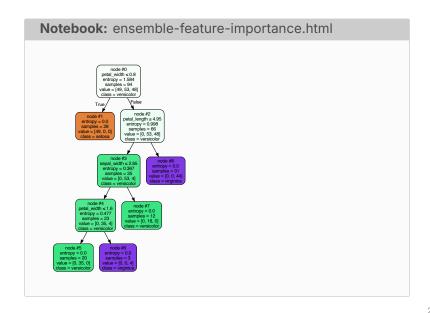


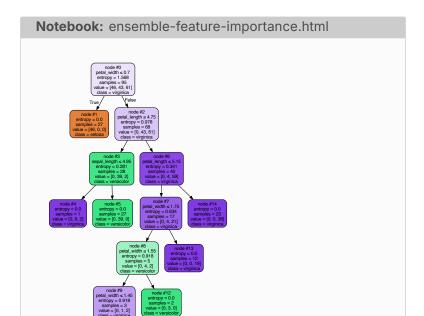


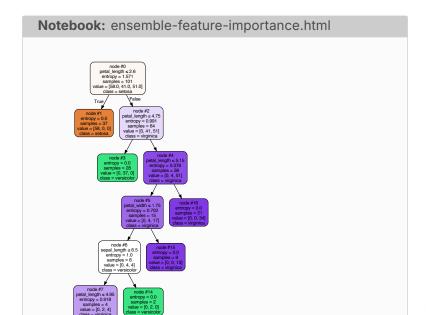


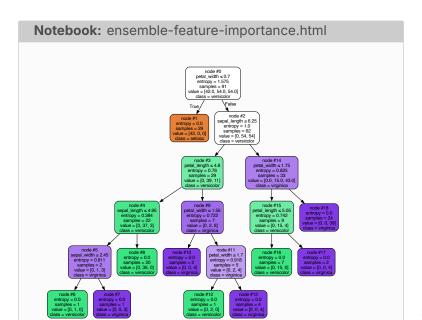


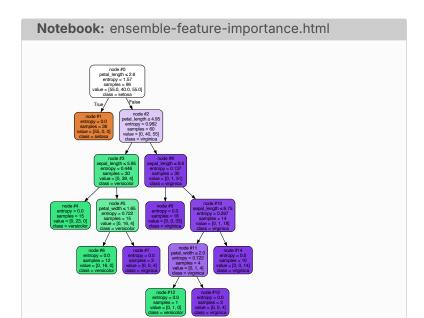




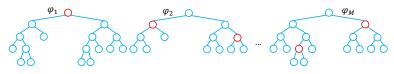








Feature Importance¹



Importance of variable X_j for an ensemble of M trees φ_m is:

$$\operatorname{Imp}(X_j) = \frac{1}{M} \sum_{m=1}^{M} \sum_{t \in \omega_m} 1(j_t = j) \Big[p(t) \Delta i(t) \Big],$$

where j_t denotes the variable used at node t, $p(t) = N_t/N$ and $\Delta i(t)$ is the impurity reduction at node t:

$$\Delta i(t) = i(t) - \frac{N_{t_L}}{N_t} i(t_L) - \frac{N_{t_r}}{N_t} i(t_R)$$

¹Slide Courtesy Gilles Louppe

Computed Feature Importance

